



MARKED-UP VERSION INDICATING CHANGES MADE

Table I

Bruker Reflex MALDI-TOF mass spectrometer
analysis of 55 kDa peptides

<u>SEQ ID</u> <u>NO</u>	Sequence	peptide mass Measured (Da)	(monoisotopic) Calculated (Da)
1	QMSLLR	859.48	859.495
2	AVDSLVPIGR	1025.58	1025.587
3	VGLKAPGIIPR	1119.68	1119.713
4	TIAMDGTTEGLVR	1261.40	1261.634
5	ISVREPMQTGIK	1357.70	1357.739
6	IMNVIGEPIDER	1384.68	1384.702
7	AHGGYSVFAGVGER	1405.66	1405.674
8	FTQAGSEVSALLGR	1434.73	1434.747
9	TSIADTIINQKR	1471.81	1471.836
10	EAYPGDVFYLSR	1552.71	1552.731
11	VALVYGQMNEPPGAR	1600.79	1600.803
12	TGAIVDPVGEELLGR	1623.87	1623.883
13	LVLEVAQHLGESTVR	1649.88	1649.910
14	IMDPNIVGSEHYDVAR	1814.85	1814.862
15	VLDSGAPIKIPVGPETLGR	1918.08	1918.089
16	AIAELGIYPAVDPLDSTS R	1986.99	1987.026
17	IMNVIGEPIDERGPIKTK	2009.10	2009.098
18	IPSAVGYQPTLATDMGTMQER	2265.06	2265.077
19	EVAAFAQFGSDLDAATQQLLSR	2337.15	2337.160